EFS

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/589,305Source: 1510Date Processed by STIC: 2/23/07

ENTERED



IFWO

RAW SEQUENCE LISTING DATE: 02/23/2007 PATENT APPLICATION: US/10/589,305 TIME: 15:04:42

Input Set : N:\efs\02_23_07\10589305_efs\SEQUENCE-LISTING. axt

Output Set: N:\CRF4\02232007\J589305.raw

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3 <110> APPLICANT: ARES TRADING S.A.
     5 <120> TITLE OF INVENTION: Leucine rich repeat containing protein
     7 <130> FILE REFERENCE: P036769WO
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/589,305
C--> 9 <141> CURRENT FILING DATE: 2006-08-14
     9 <150> PRIOR APPLICATION NUMBER: GB 0403142.3
    10 <151> PRIOR FILING DATE: 2004-02-12
    12 <160> NUMBER OF SEQ ID NOS: 50
    14 <170> SOFTWARE: SeqWin99, version 1.02
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    17 <211> LENGTH: 496
    18 <212> TYPE: DNA
    19 <213> ORGANISM: Homo sapiens
    21 <400> SEQUENCE: 1
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    23 atgacaaatg catcaaqaaa aagcaatatt ttattcaatt ctgaatgcca atggaatgaa
                                                                             120
    24 tatattctga caaattgttc ttttaccgga aagtgtgata tacctgtgga catatcacag
                                                                             180
    25 acagcagcca ctgtggatgt aagtttcaat ttctttagag ttctcttaca gtctcacacg
    26 aaaaaaqaag agtggaaaat aaaacatctg gacctcagta acaatctcat atcaaaaata
                                                                             300
    27 accttaagcc cttttgcata tttacatgct ttggaagtgt taaacctcag caacaatgcc
                                                                             360
    28 atccactccc tctcattgga tctactcagt cctaagtcct catgggtgaa acgccacaga
                                                                             420
    29 agcagettea gaaacaggtt tecattgetg aaggtgetea ttetteaaag aaataaaete
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                                                                             496
    30 agtgacactc ccaagg
    32 <210> SEQ ID NO: 2
    33 <211> LENGTH: 166
    34 <212> TYPE: PRT
    35 <213> ORGANISM: Homo sapiens
    37 <400> SEQUENCE: 2
    38 Met Lys Asn Leu Tyr Phe Arg Val Ile Thr Ile Val Ile Gly Leu Tyr
    39 1
                       5
    41 Phe Thr Gly Ile Met Thr Asn Ala Ser Arg Lys Ser Asn Ile Leu Phe
    42
    44 Asn Ser Glu Cys Gln Trp Asn Glu Tyr Ile Leu Thr Asn Cys Ser Phe
    45
                                    40
    47 Thr Gly Lys Cys Asp Ile Pro Val Asp Ile Ser Gln Thr Ala Ala Thr
    48
                                55
    50 Val Asp Val Ser Phe Asn Phe Phe Arg Val Leu Leu Gln Ser His Thr
    53 Lys Lys Glu Glu Trp Lys Ile Lys His Leu Asp Leu Ser Asn Asn Leu
                        85
                                            90
    56 Ile Ser Lys Ile Thr Leu Ser Pro Phe Ala Tyr Leu His Ala Leu Glu
                   100
                                        105
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59 Val Leu Asn Leu Ser Asn Asn Ala Ile His Ser Leu Ser Leu Asp Leu

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                               120
62 Leu Ser Pro Lys Ser Ser Trp Val Lys Arg His Arg Ser Ser Phe Arg
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65 Asn Arg Phe Pro Leu Leu Lys Val Leu Ile Leu Gln Arg Asn Lys Leu
                                            155
66 145
                       150
68 Ser Asp Thr Pro Lys Gly
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72 <211> LENGTH: 170
73 <212> TYPE: DNA
74 <213 > ORGANISM: Homo sapiens
76 <400> SEQUENCE: 3
77 qactqtqqaa actqaaqtca ttqcaqaqtt tqqatctqtc attcaatqqq atattqcaaa
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78 tagggtggtc tgattttcac aactgcctgc aactggagaa tctctgttta aagagcaaca
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79 agatattcaa aattccccca caagccttca aggacctcaa aaaattacag
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81 <210> SEQ ID NO: 4
82 <211> LENGTH: 56
83 <212> TYPE: PRT
84 <213> QRGANISM: Homo sapiens
86 <400> SEQUENCE: 4
87 Leu Trp Lys Leu Lys Ser Leu Gln Ser Leu Asp Leu Ser Phe Asn Gly
90 Ile Leu Gln Ile Gly Trp Ser Asp Phe His Asn Cys Leu Gln Leu Glu
               20
                                   25
93 Asn Leu Cys Leu Lys Ser Asn Lys Ile Phe Lys Ile Pro Pro Gln Ala
                                                    45
94
           35
96 Phe Lys Asp Leu Lys Lys Leu Gln
       50
99 <210> SEQ ID NO: 5
100 <211> LENGTH: 190
101 <212> TYPE: DNA
102 <213> ORGANISM: Homo sapiens
104 <400> SEQUENCE: 5
105 gtcatagacc ttagcaacaa tgctctgatt accatcctac caatgatgat catagctcta
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106 qaatttcccc atctagtggt tgacttggct gataataact ggcagtgtga tgatagtgtg
                                                                          120
107 gcagtettte aaaattttat ttetgaatee tggaggaaaa agtggaatgt catttgcaae
                                                                          180
                                                                          190
108 aggtctatag
110 <210> SEQ ID NO: 6
111 <211> LENGTH: 64
112 <212> TYPE: PRT
113 <213> ORGANISM: Homo sapiens
115 <400> SEQUENCE: 6
116 Val Ile Asp Leu Ser Asn Asn Ala Leu Ile Thr Ile Leu Pro Met Met
117 1
119 Ile Ile Ala Leu Glu Phe Pro His Leu Val Val Asp Leu Ala Asp Asn
                20
122 Asn Trp Gln Cys Asp Asp Ser Val Ala Val Phe Gln Asn Phe Ile Ser
                                40
125 Glu Ser Trp Arg Lys Lys Trp Asn Val Ile Cys Asn Arg Ser Ile Gly
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129 <210> SEQ ID NO: 7
130 <211> LENGTH: 1754
131 <212> TYPE: DNA
132 <213> ORGANISM: Homo sapiens
134 <400> SEQUENCE: 7
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136 ctcccattca tctgcatcgc atgaaaagcc tcataaggag caaagcagag aggccccagg
                                                                         120
137 gaggaaggca cacgggcatt tctactctgg ggaagaaggc aaaggccggc tctggtctca
                                                                         180
138 ggaagaagca gagacggctg ccaaggagtg ttagaagcac ccgcgatgtg caggctgccg
                                                                         240
139 gcaaaaaaga ggacgctccc caggacctgg ctctggcggt gtgcctgtca gtgttcatca
                                                                         300
140 catteettgt egeetteage etgggggett teacaaggee ttatgttgae agaetgtgge
                                                                         360
141 aaaaaaagtg ccagagcaaa agccctggcc tggacaacgc gtattcaaac gagggcttct
                                                                         420
142 acgatgacat ggaagctgcg gggcacacac cacacccaga gacccatctg cgccaagtat
                                                                         480
143 tteeteatet aageetetae gagaaceaga eeeetttetg ggtgacacag eeacaceeae
                                                                         540
144 acgccaccgt aattcctgat agaactctgg gaaggagcag aaaggatcct ggcagttcgc
                                                                         600
145 agagcccagg acagtgcggg gacaacaccg gggcaggaag tggaaatgat ggtgcagtct
                                                                         660
146 attecattet ceagagaeat ceacatgeeg gtaacegtga actaatgtea geagegeagg
                                                                         720
147 accacateca taggaatgat atteteggag aatggaetta tgaaaetgtg geecaggaag
                                                                         7.80
148 agecteteag tgeacattea gtgggegtet ettetgtage tggeacgtet caegetgtet
                                                                         '840
149 ctggctcaag ccgttatgat tccaatgaat tagacccttc cctctccgga gaaataacag
                                                                         900
150 cttccctctg taaaatgcta acacatgcag aagcacagag gactggagat agtaaggaaa
                                                                         960
151 gagggggcac tgaacagtca ctttgggact cgcagatgga attttctaag gaaaggcaag
                                                                         1020
152 tgagttcatc cattgatttg ctgagcatac agcagccaag gctgtccggg gcaagggctg
                                                                         1080
153 aggaageget tteageecae tacagegagg ttecataegg tgacecaaga gacacaggee
                                                                         1140
154 catcagtett tectecaaga tgggacagtg geetggatgt caeteetget aacaaggaac
                                                                         1200
155 cagtgcagaa atccactcct tctgacactt gctgtgagtt ggagagtgac tgtgactctg
                                                                         1260
156 atgaggggtc tctgttcact ctgagctcca taagttcaga gagtgcaagg agcaagactg
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157 aagaggcagt gcctgatgag gagtccctgc aggacgagag ctcaggggca agcaaggaca
158 atgtgacggc tgtagacagt cttgaggaaa atgttacctt ccaaacaatt ccagggaaat
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159 gcaagaatca agaagatccc tttgaaaaac ctctcatttc tgctccagac tctggcatgt
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160 acaagactca tctggaaaat gcctctgaca ctgatagatc tgagggcctg tcaccctggc
                                                                         1560
161 ccaggtcacc agggaatagt cccttagggg atgagtttcc gggcatgttc acttatgatt
                                                                         1620
162 atgacacage tetteaatee aaggeageag aatggeattg eteaettaga gaettagaat
                                                                         1680
163 tttcaaatgt ggacgtttta cagcaaacac caccatgttc tgctgaagtt ccctcagatc
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164 ctgataaggc tgcc
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166 <210> SEO ID NO: 8
167 <211> LENGTH: 584
168 <212> TYPE: PRT
169 <213> ORGANISM: Homo sapiens
171 <400> SEQUENCE: 8
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173 1
175 Thr Arg Leu Pro Pro Ile His Leu His Arg Met Lys Ser Leu Ile Arg
176
                20
178 Ser Lys Ala Glu Arg Pro Gln Gly Gly Arg His Thr Gly Ile Ser Thr
181 Leu Gly Lys Lys Ala Lys Ala Gly Ser Gly Leu Arg Lys Lys Gln Arg
182
                            55
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184 Arg Leu Pro Arg Ser Val Arg Ser Thr Arg Asp Val Gln Ala Ala Gly 185 65 187 Lys Lys Glu Asp Ala Pro Gln Asp Leu Ala Leu Ala Val Cys Leu Ser 190 Val Phe Ile Thr Phe Leu Val Ala Phe Ser Leu Gly Ala Phe Thr Arg 191 100 105 193 Pro Tyr Val Asp Arg Leu Trp Gln Lys Lys Cys Gln Ser Lys Ser Pro 115 120 196 Gly Leu Asp Asn Ala Tyr Ser Asn Glu Gly Phe Tyr Asp Asp Met Glu 130 135 199 Ala Ala Gly His Thr Pro His Pro Glu Thr His Leu Arg Gln Val Phe 200 145 150 155 202 Pro His Leu Ser Leu Tyr Glu Asn Gln Thr Pro Phe Trp Val Thr Gln 165 170 205 Pro His Pro His Ala Thr Val Ile Pro Asp Arg Thr Leu Gly Arg Ser 180 185 208 Arg Lys Asp Pro Gly Ser Ser Gln Ser Pro Gly Gln Cys Gly Asp Asn 195 200 211 Thr Gly Ala Gly Ser Gly Asn Asp Gly Ala Val Tyr Ser Ile Leu Gln 215 220 214 Arg His Pro His Ala Gly Asn Arg Glu Leu Met Ser Ala Ala Gln Asp 230 235 217 His Ile His Arg Asn Asp Ile Leu Gly Glu Trp Thr Tyr Glu Thr Val 245 250 220 Ala Gln Glu Glu Pro Leu Ser Ala His Ser Val Gly Val Ser Ser Val 260 265 270 223 Ala Gly Thr Ser His Ala Val Ser Gly Ser Ser Arg Tyr Asp Ser Asn 275 280 226 Glu Leu Asp Pro Ser Leu Ser Gly Glu Ile Thr Ala Ser Leu Cys Lys 295 229 Met Leu Thr His Ala Glu Ala Gln Arg Thr Gly Asp Ser Lys Glu Arg 230 305 232 Gly Gly Thr Glu Gln Ser Leu Trp Asp Ser Gln Met Glu Phe Ser Lys 325 330 235 Glu Arg Gln Val Ser Ser Ser Ile Asp Leu Leu Ser Ile Gln Gln Pro 340 345 238 Arg Leu Ser Gly Ala Arg Ala Glu Glu Ala Leu Ser Ala His Tyr Ser 355 360 365 241 Glu Val Pro Tyr Gly Asp Pro Arg Asp Thr Gly Pro Ser Val Phe Pro 375 380 244 Pro Arg Trp Asp Ser Gly Leu Asp Val Thr Pro Ala Asn Lys Glu Pro 395 390 . 247 Val Gln Lys Ser Thr Pro Ser Asp Thr Cys Cys Glu Leu Glu Ser Asp 405 410 250 Cys Asp Ser Asp Glu Gly Ser Leu Phe Thr Leu Ser Ser Ile Ser Ser 251 425 253 Glu Ser Ala Arg Ser Lys Thr Glu Glu Ala Val Pro Asp Glu Glu Ser 440 256 Leu Gln Asp Glu Ser Ser Gly Ala Ser Lys Asp Asn Val Thr Ala Val

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257
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                             455
259 Asp Ser Leu Glu Glu Asn Val Thr Phe Gln Thr Ile Pro Gly Lys Cys
260 465
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262 Lys Asn Gln Glu Asp Pro Phe Glu Lys Pro Leu Ile Ser Ala Pro Asp
263
                    485
                                         490
265 Ser Gly Met Tyr Lys Thr His Leu Glu Asn Ala Ser Asp Thr Asp Arg
                500
                                     505
268 Ser Glu Gly Len Ser Pro Trp Pro Arg Ser Pro Gly Asn Ser Pro Leu
269
            515
                                520
                                                     525
271 Gly Asp Glu Phe Pro Gly Met Phe Thr Tyr Asp Tyr Asp Thr Ala Leu
        530
                            535
274 Gln Ser Lys Ala Ala Glu Trp His Cys Ser Leu Arg Asp Leu Glu Phe
275 545
                        550
                                             555
277 Ser Asn Val Asp Val Leu Gln Gln Thr Pro Pro Cys Ser Ala Glu Val
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                                         570
280 Pro Ser Asp Pro Asp Lys Ala Ala
281
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283 <210> SEQ ID NO: 9
284 <211> LENGTH: 2610
285 <212> TYPE: DNA
286 <213> ORGANISM: Homo sapiens
288 <400> SEQUENCE: 9
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290 atgacaaatg catcaagaaa aagcaatatt ttattcaatt ctgaatgcca atggaatgaa
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291 tatattetga caaattgtte ttttaeegga aagtgtgata taeetgtgga catateaeag
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292 acagcagcca ctgtggatgt aagtttcaat ttctttagag ttctcttaca gtctcacacg
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293 aaaaaagaag agtggaaaat aaaacatctg gacctcagta acaatctcat atcaaaaata
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294 accttaagcc cttttgcata tttacatgct ttggaagtgt taaacctcag caacaatgcc
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295 atccactccc tctcattgga tctactcagt cctaagtcct catgggtgaa acgccacaga
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296 agcagettea gaaacaggtt tecattgetg aaggtgetea ttetteaaag aaataaaete
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297 agtgacactc ccaagggact gtggaaactg aagtcattgc agagtttgga tctgtcattc
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298 aatgggatat tgcaaatagg gtggtctgat tttcacaact gcctgcaact ggagaatctc
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299 tgtttaaaga gcaacaagat attcaaaatt cccccacaag ccttcaagga cctcaaaaaa
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300 ttacaggtca tagaccttag caacaatgct ctgattacca tcctaccaat gatgatcata
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301 getetagaat tteeceatet agtggttgae ttggetgata ataactggea gtgtgatgat
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302 agtgtggcag tctttcaaaa ttttatttct qaatcctgga ggaaaaagtg gaatgtcatt
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303 tgcaacaggt ctatagggag tgaggaggcc aacgggggca ctccccagag caggatttcc
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304 agggaaaccc gccttcctcc cattcatctg catcgcatga aaagcctcat aaggagcaaa
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305 gcagagaggc cccagggagg aaggcacacg ggcatttcta ctctggggaa gaaggcaaag
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306 gccggctctg gtctcaggaa gaagcagaga cggctgccaa ggagtgttag aagcacccgc
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307 gatgtgcagg ctgccggcaa aaaagaggac gctccccagg acctggctct ggcggtgtgc
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308 ctgtcagtgt tcatcacatt ccttgtcgcc ttcagcctgg gggctttcac aaggccttat
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309 gttgacagac tgtggcaaaa aaagtgccag agcaaaagcc ctggcctgga caacgcgtat
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310 tcaaacgagg gcttctacga tgacatggaa gctgcggggc acacaccaca cccagagacc
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311 catctgegec aagtatttee teatetaage etetaegaga accagacece tttetgggtg
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312 acacagecae acecacaege cacegtaatt cetgatagaa etetgggaag gageagaaag
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313 gatcctggca gttcgcagag cccaggacag tgcggggaca acaccggggc aggaagtgga
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314 aatgatggtg cagtctattc cattctccag agacatccac atgccggtaa ccgtgaacta
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315 atgtcagcag cgcaggacca catccatagg aatgatattc tcggagaatg gacttatgaa
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VERIFICATION SUMMARY PATENT APPLICATION: US/10/589,305

DATE: 02/23/2007 TIME: 15:04:43

Input Set : N:\efs\02_23_07\10589305_efs\SEQUENCE-LISTING.txt
Output Set: N:\CRF4\02232007\J589305.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date